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(54) Title: METHOD FOR DETECTING AND QUANTIFYING RARE MUTATIONS/POLYMORPHISMS

(57) Abstract: The present invention is directed to a method for detecting and quantifying rare mutations in a nucleic acid sample. The nucleic acid molecules under investigation can be either DNA or RNA. The rare mutation can be any type of functional or non-functional nucleic acid change or mutation, such as deletion, insertion, translocation, inversion, one or more base substitution or polymorphism. Therefore, the methods of the present invention are useful in detection of rare mutations in, for example, diagnostic, prognostic and follow-up applications, when the targets are rare known nucleic acid variants mixed in with the wildtype or the more common nucleic acid variant(s).



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